

M C P S Q P T Y P G D P G P
ATG TGC CCG AGC CAG CCG ACC TAT CCG GGC GAT CCC GGG CCG
V E D L I R F Y D N L Q Q W
GTG GAA GAT CTG ATC CGC TTT TAT GAT AAC CTG CAG CAG TGG
L N C V T A A C *
CTG AAC TGC GTG ACC GCC GCC TGC TAG

FIGURE 1

1 11 21 31 41
ACACACCATA TGTCCCCGAG CCAGCCGACC TATCCGGGC CG ATCCCGGGCC
TGTGTGGTAT ACACGGGCTC GGTGGCTGG ATAGGCCCGC TAGGGCCCGG

51 61 71 81 91
GGTGAAGAT CTGATCCGCT TTATGATAA CCTGCAGCAG TGGCTGAAC T
CCACCTTCTA GACTAGGC GA AAATACTATT GGACGTCGTC ACCGACTTGA

101 111 121 131
GCGTGACCGC CGCCTGCTAG GGATCCACAC AC
CGCACTGGCG GCGGACGATC CCTAGGTGTG TG

FIGURE 2

10027033 * 122004



FIGURE 3

10027038 - 122001

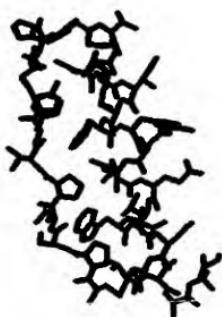


FIGURE 4

10027038, 122001



FIGURE 5

10027038.122064

PRO227-8502200T

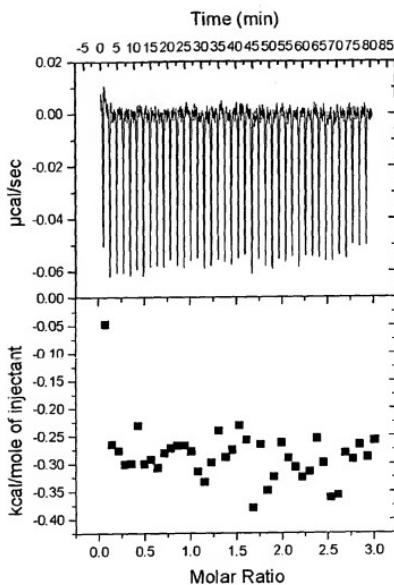


FIGURE 6

T0022T = 3202300T

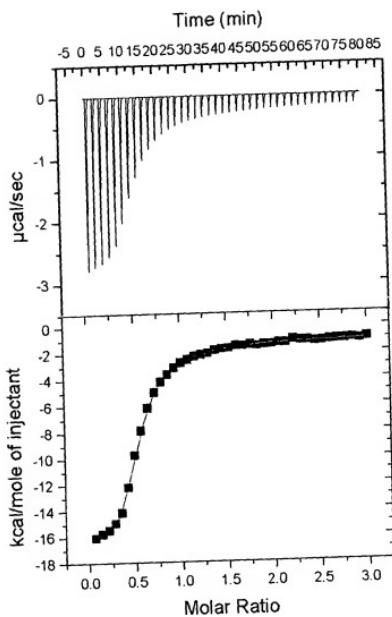


FIGURE 7

400027038 "422000

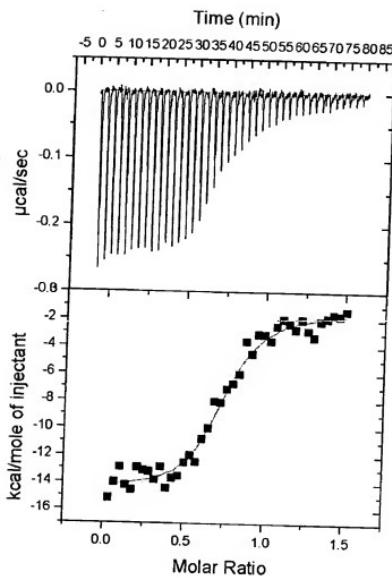


FIGURE 8

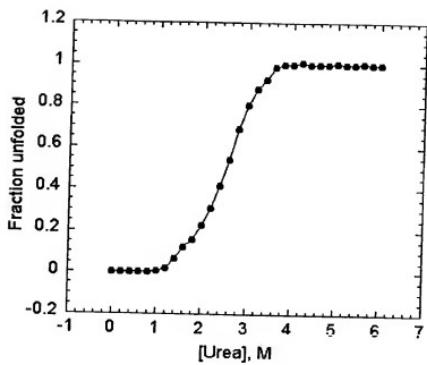


FIGURE 9

10627038 - 122004

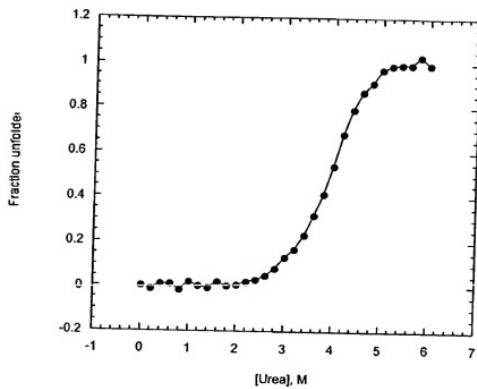


FIGURE 10

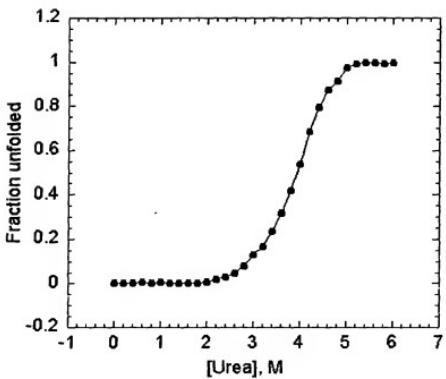


FIGURE 11

DOI:10.2210/aob/m/2008/100

Y0022T "85022003"

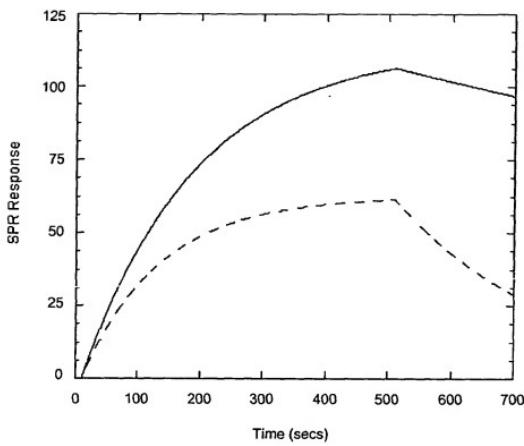


FIGURE 12

10227038 - 422081

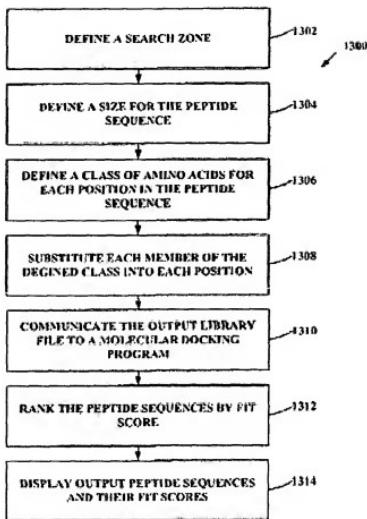


FIG. 13

40022038 402203
14

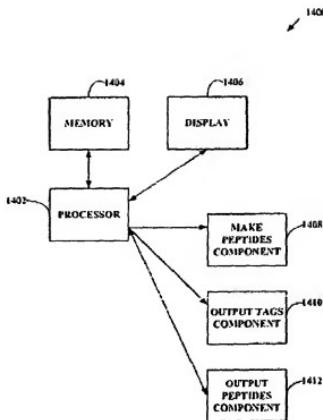


FIG. 14

1002038-1-1
Jb

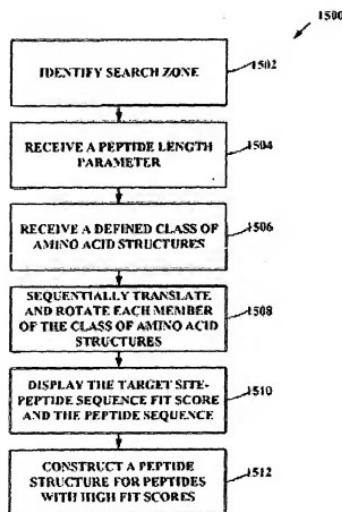


FIG. 15